



#5

SEQUENCE LISTING

<110> Afar, Daniel  
Hubert, Rene S.  
Leong, Kahan  
Raitano, Arthur B.  
Saffran, Douglas C.  
Mitchell, Steve Chappell

<120> PEPTIDES DERIVED FROM STEAP1 (AS AMENDED)

<130> 511582001601

<140> US 10/010,667  
<141> 2001-12-06

<150> 09/323,873  
<151> 1999-06-01

<150> 60/087,520  
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<150> 60/091,183  
<151> 1998-06-30

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<220>  
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Gly Glu Thr Ser Met Leu Lys Arg Pro Val Leu Leu His Leu His Gln  
35 40 45  
Thr Ala His Ala Asp Glu Phe Asp Cys Pro Ser Glu Leu Gln His Thr  
50 55 60  
Gln Glu Leu Phe Pro Gln Trp His Leu Pro Ile Lys Ile Ala Ala Ile  
65 70 75 80  
Ile Ala Ser Leu Thr Phe Leu Tyr Thr Leu Leu Arg Glu Val Ile His  
85 90 95  
Pro Leu Ala Thr Ser His Gln Gln Tyr Phe Tyr Lys Ile Pro Ile Leu  
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115 120 125  
Val Tyr Leu Pro Gly Val Ile Ala Ala Ile Val Gln Leu His Asn Gly  
130 135 140  
Thr Lys Tyr Lys Phe Pro His Trp Leu Asp Lys Trp Met Leu Thr  
145 150 155 160  
Arg Lys Gln Phe Gly Leu Leu Ser Phe Phe Ala Val Leu His Ala  
165 170 175  
Ile Tyr Ser Leu Ser Tyr Pro Met Arg Arg Ser Tyr Arg Tyr Lys Leu  
180 185 190  
Leu Asn Trp Ala Tyr Gln Gln Val Gln Gln Asn Lys Glu Asp Ala Trp  
195 200 205  
Ile Glu His Asp Val Trp Arg Met Glu Ile Tyr Val Ser Leu Gly Ile  
210 215 220  
Val Gly Leu Ala Ile Leu Ala Leu Leu Ala Val Thr Ser Ile Pro Ser  
225 230 235 240  
Val Ser Asp Ser Leu Thr Trp Arg Glu Phe His Tyr Ile Gln Ser Lys  
245 250 255  
Leu Gly Ile Val Ser Leu Leu Leu Gly Thr Ile His Ala Leu Ile Phe  
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Ala Trp Asn Lys Trp Ile Asp Ile Lys Gln Phe Val Trp Tyr Thr Pro  
275 280 285  
Pro Thr Phe Met Ile Ala Val Phe Leu Pro Ile Val Val Leu Ile Phe  
290 295 300  
Lys Ser Ile Leu Phe Leu Pro Cys Leu Arg Lys Lys Ile Leu Lys Ile  
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ggattactaa	gtttttctt	cgctatggtc	catgttgcct	acaggcttg	cttaccgatg	240
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ttaaactgg	gagaatttcg	tttatttcg	tctacacttg	gatatgtcg	tctgtctata	480
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<213> Homo sapiens

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Ala Ala Tyr Gln Leu Tyr Tyr Gly Thr Lys Tyr Arg Arg Phe Pro Pro  
35 40 45  
Trp Leu Glu Thr Trp Leu Gln Cys Arg Lys Gln Leu Gly Leu Leu Ser  
50 55 60  
Phe Phe Phe Ala Met Val His Val Ala Tyr Ser Leu Cys Leu Pro Met  
65 70 75 80  
Arg Arg Ser Glu Arg Tyr Leu Phe Leu Asn Met Ala Tyr Gln Gln Val  
85 90 95  
His Ala Asn Ile Glu Asn Ser Trp Asn Glu Glu Glu Val Trp Arg Ile  
100 105 110  
Glu Met Tyr Ile Ser Phe Gly Ile Met Ser Leu Gly Leu Leu Ser Leu  
115 120 125  
Leu Ala Val Thr Ser Ile Pro Ser Val Ser Asn Ala Leu Asn Trp Arg  
130 135 140  
Glu Phe Ser Phe Ile Gln Ser Thr Leu Gly Tyr Val Ala Leu Leu Ile  
145 150 155 160  
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165 170

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<212> DNA  
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attactttgc tctccctagt atacccgtca ggtcttcgg cagctgctta tcaactttat 180  
tacggcacca agtataggag atttccacct tggtggaaa cctggttaca gtgtggaaaa 240  
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<210> 11  
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atCTTGTGTA CAGCCCACAC CCTGGGTGAC ggtGGGAAGA gATTCTCAG CCCTTAAAT	240	
CTCAGATGGT ATCTTCCCTGC AGCCTACGTG ttAGGGCTTA TCATTCCTG CACTGTGCTG	300	
gtGATCAgT TTGTCTTAAT CATGCCATGT gtAGACAACA CCCTTACAAG gATCCGCCAG	360	
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gCTCAACTGG AGGGAGTTCA GCTTCGTCA GTCCTACTG GGCTTGTGG CCNTCGTGT	240	
gAGCACACtN CACACGCTCA CCTACGGGTG GACCGCGCC TtCGAGGAGA GCCGCTACAA	300	
gttCTACtN CCTCCACtN tCACGNTCAc GCTGCTGGTG CCCTGCGTtC gttCATCCTG	360	
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<220>  
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<213> Homo sapiens

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